

000000000000000000000000

HU : MGSDVRDLNALLPAVPSLGGGGCALPVSGAAQWAPVLDFAAPPGASAYGSL
MO : MGSDVRDLNALLPAVSSLGGGGCGLPVSGAAQWAPVLDFAAPPGASAYGSL

HU : GGPAPPAPPAPPAPPAPPHSFIKQEPMWGAEPEEQCLSAFTVHFSGQFTGTAG
MO : GGPAPPAPPAPPAPPAPPHSFIKQEPMWGAEPEEQCLSAFTLHFSGQFTGTAG

HU : ACRYGPFGPPPSQASSGQARMFPNAPYLPSCLESQPAIRNQGYSTVTFDGTPS
MO : ACRYGPFGPPPSQASSGQARMFPNAPYLPSCLESQPTIRNQGYSTVTFDGAPS

HU : YGHTPSHAAQFPNHSFKEDPMGQQGSLGEQQYSVPPPVGCHTPTDCTG
MO : YGHTPSHAAQFPNHSFKEDPMGQQGSLGEQQYSVPPPVGCHTPTDCTG

HU : SQALLLRTPYSSDNLYQMITSQLECMWNQMNLGATLKGVAAGSSSSVKWTE
MO : SQALLLRTPYSSDNLYQMITSQLECMWNQMNLGATLKGMAAGSSSSVKWTE

HU : GQSNHSTGYESDNHTTPILCGAQYRIHTHGVFRGIQDVRRVPGVAPTLVRSAS
MO : GQSNHGIGYESDNHTAPILCGAQYRIHTHGVFRGIQDVRRVSGVAPTLVRSAS

HU : ETSEKRPFMCAYPGCNKRYFKLSHLQMHSRKHTGEKPYQCDFKDCERRFSR
MO : ETSEKRPFMCAYPGCNKRYFKLSHLQMHSRKHTGEKPYQCDFKDCERRFSR

HU : SDQLKRHQRRHTGVKPFQCKTCQRKFSSRSDHLKTHTRHTGKTSEKPFSCR
MO : SDQLKRHQRRHTGVKPFQCKTCQRKFSSRSDHLKTHTRHTGKTSEKPFSCR

HU : WPSCQKKFARSDELVRHHNMHQRNMTKLQLAL
MO : WHSCQKKFARSDELVRHHNMHQRNMTKLHVAL

FIG. 1

860360 - 2249160

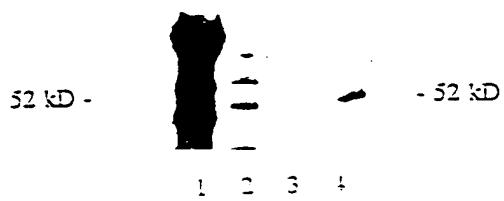


FIG. 2

0 0 0 0 0 0 0 0 0 0

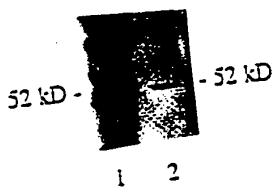
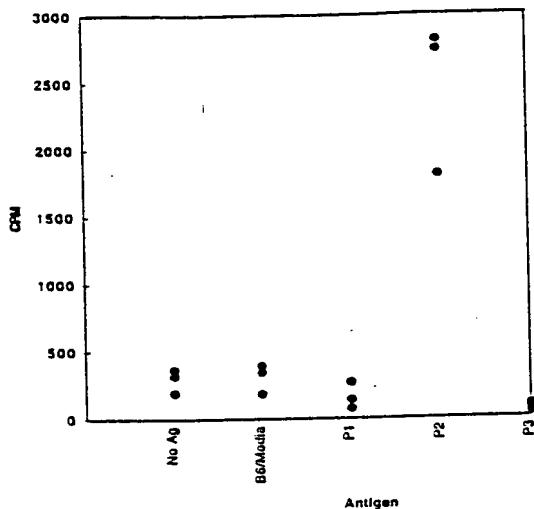


FIG. 3

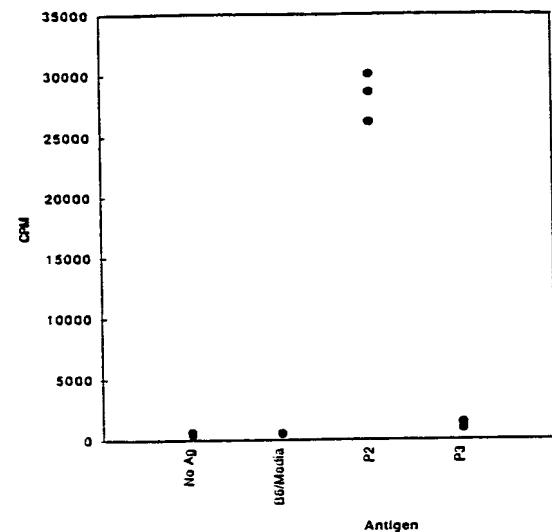


FIG. 4

A



B



C

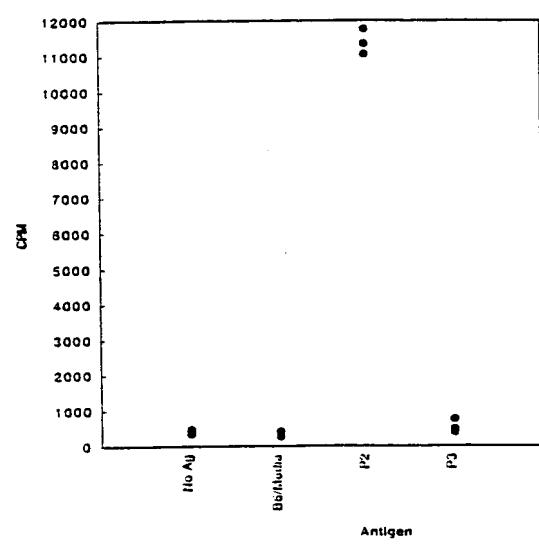
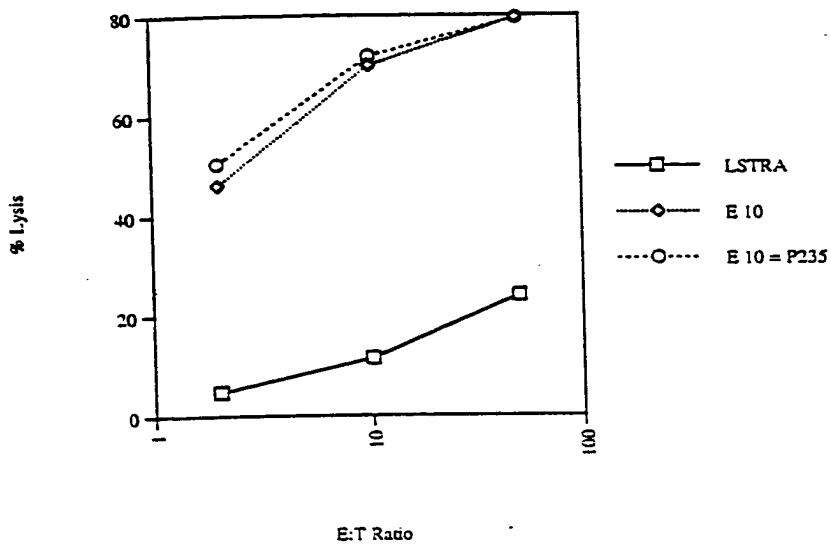


FIG. 5A-5C

A



B

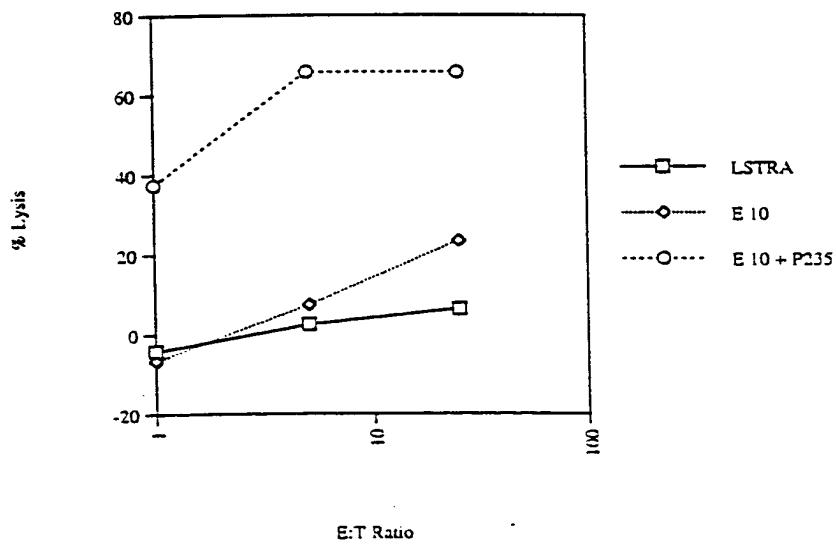


FIG. 6A and 6B

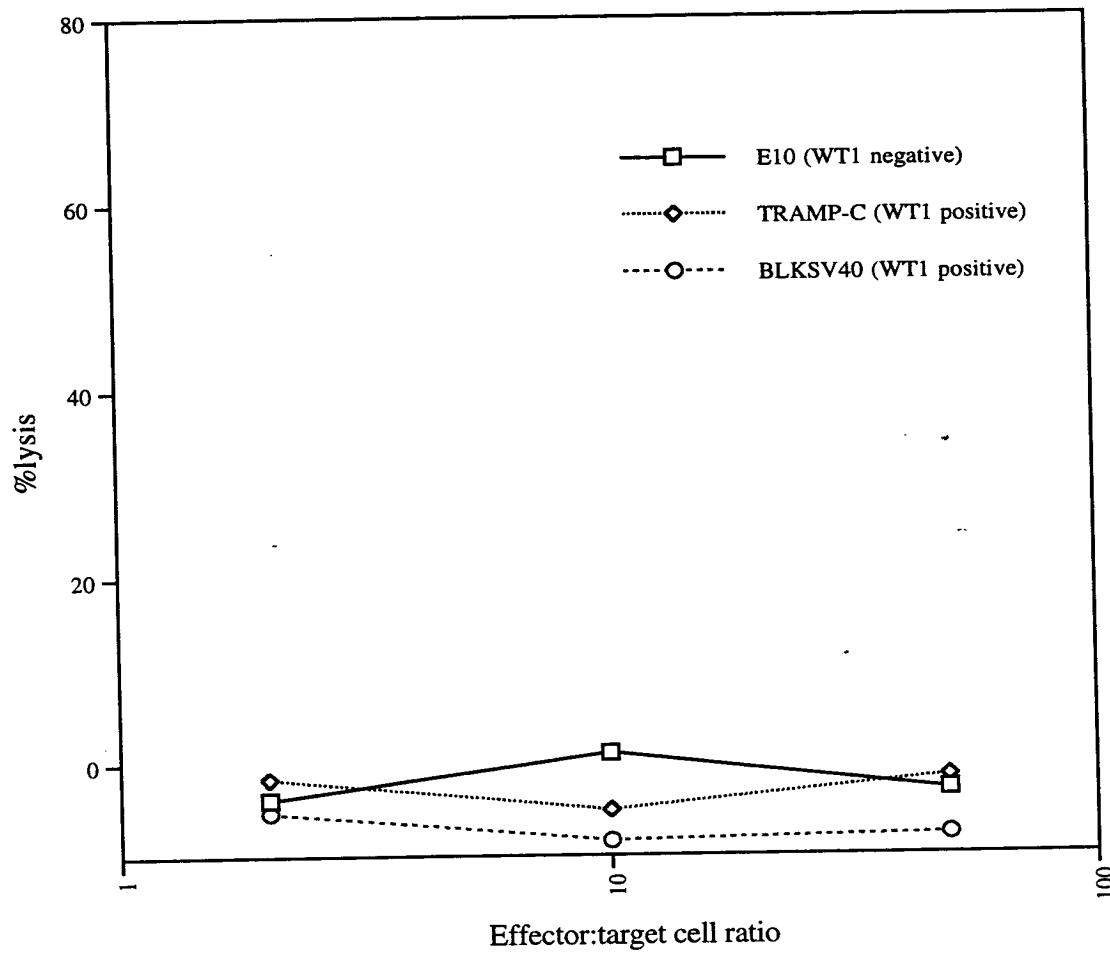


FIG. 7A

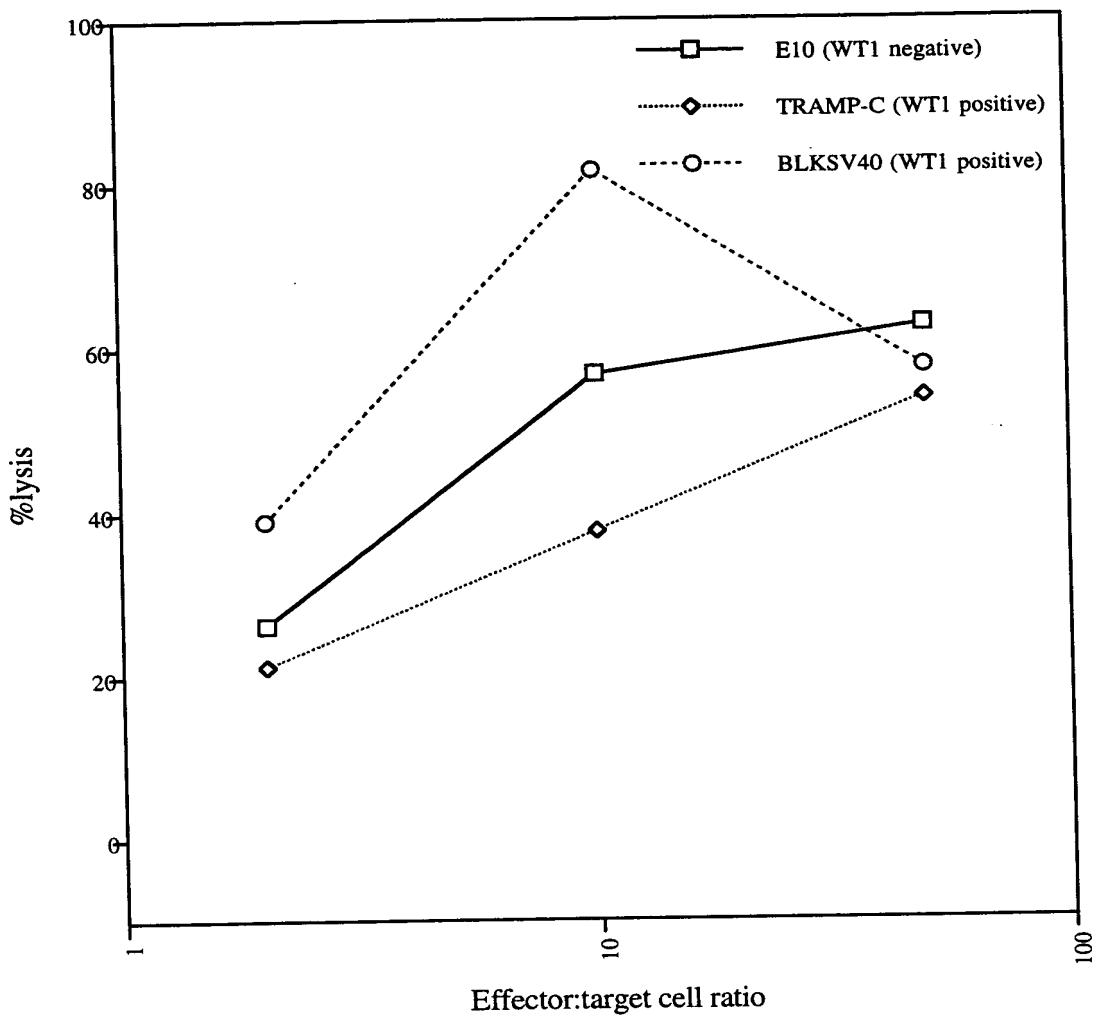


FIG. 7B

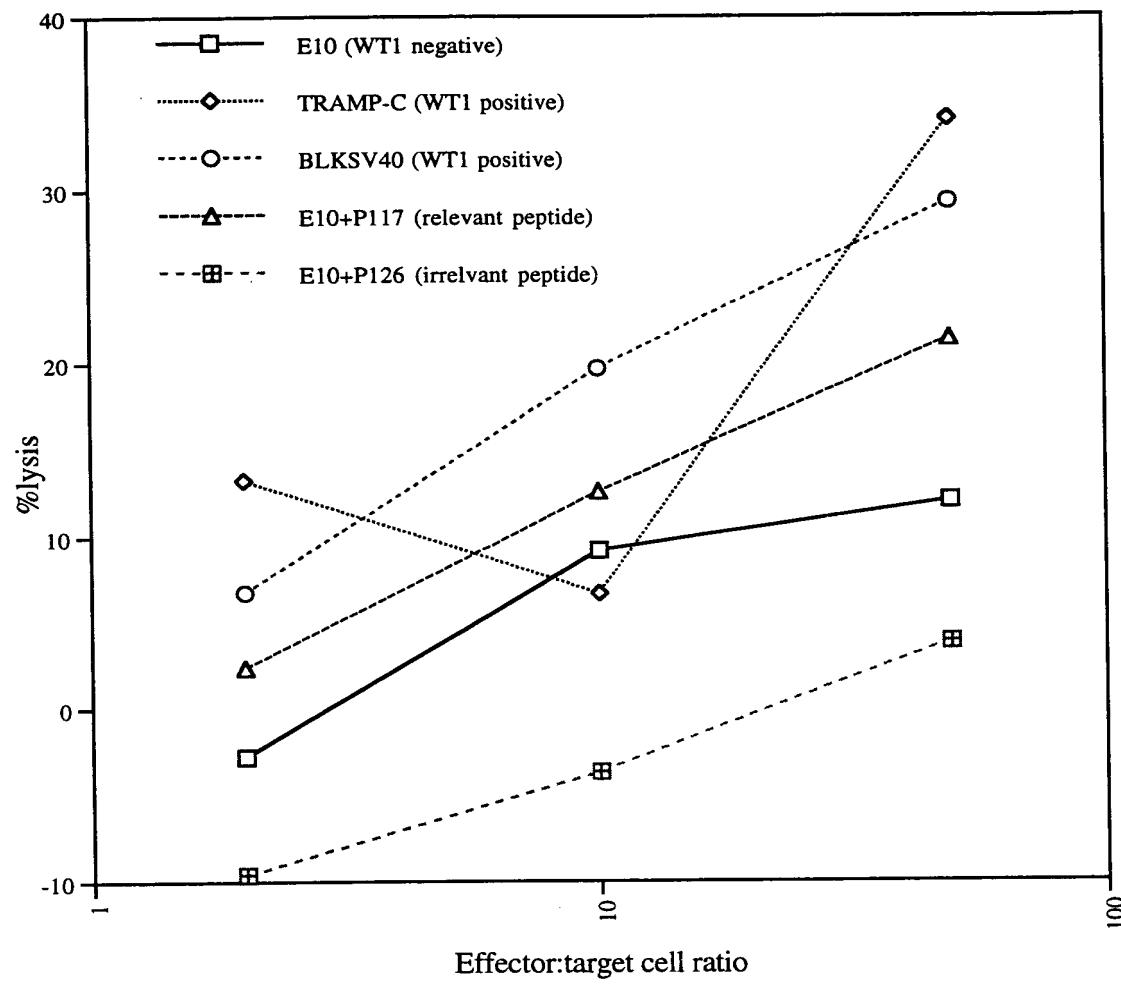


FIG. 7C

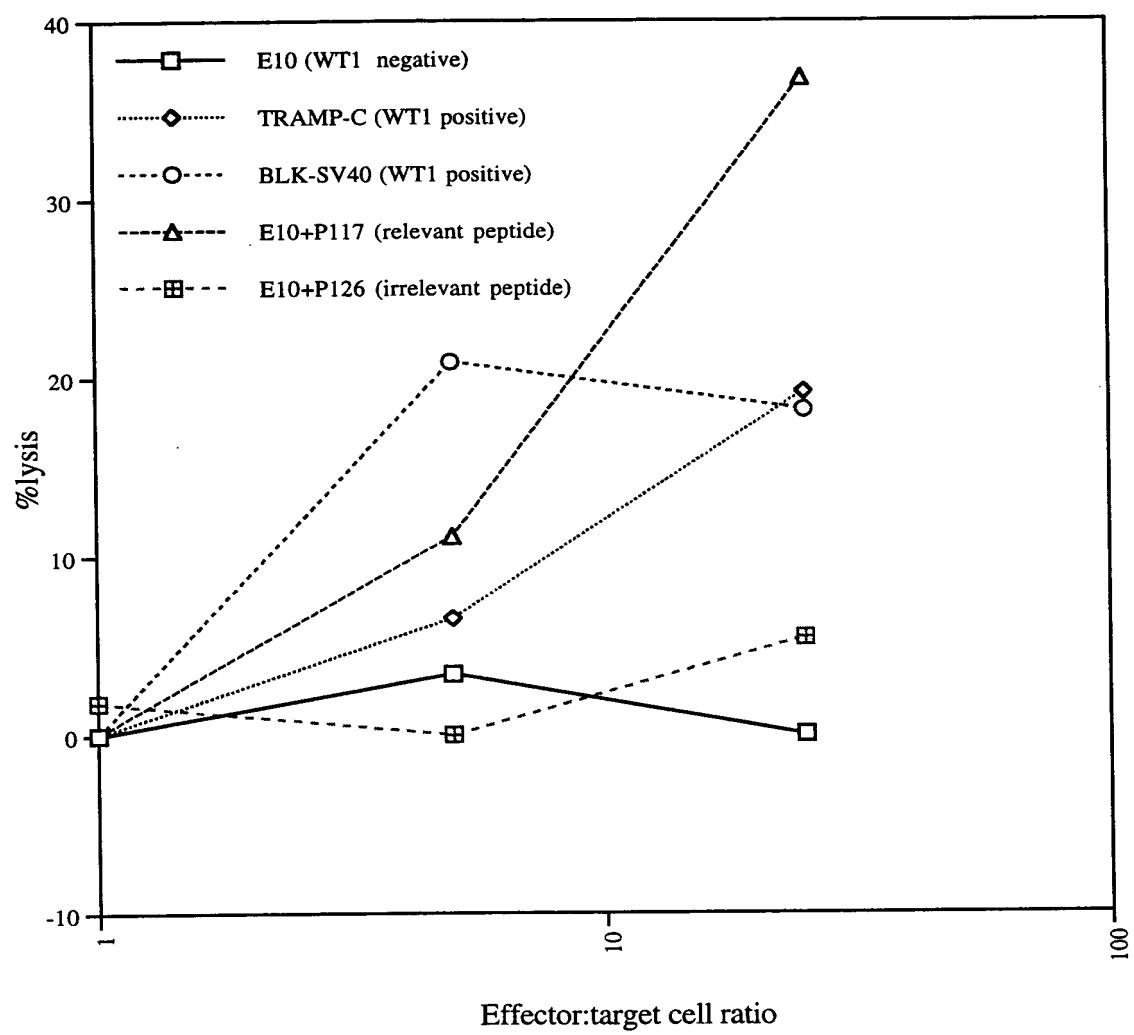


FIG. 7D

FIG. 8A

5 10 15 20 25 30 35 40 45 50 55 60 65 70 75
MGSDVRDLNALLPAVSSLGGGGCGLPVSGAAQWAPVLDFAAPPGASAYGSGLGGPAPPPAPPPPPPPPHSFIKQE
.....AAAAAAA.....AAAAA.....AAAAA.....AAAAA.....
.....RRRR.....

80 85 90 95 100 105 110 115 120 125 130 135 140 145 150
PSWGGAEPHEEQCLSAFTLHFSGQFTGTAGACRYGPFGPPPSQASSGQARMFPNAPYLPSCLESQPTIRNQGYSP
.....AAAA.....AAA.....AAAAA.....
.....RRRR.....RRRR.....
.....DDDDDDDD.....

155 160 165 170 175 180 185 190 195 200 205 210 215 220 225
TVTFDGAPSYGHTPSHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPVGCHTPDSCTGSQALLRTPYSSDN
.....AAAAA.....AAAAA.....AA
.....RRRR.....
.....DDDDDDDDDDDDDD.....

230 235 240 245 250 255 260 265 270 275 280 285 290 295 300
LYQMITSQLECMTNQMNLGATIKGMAAGSSSVKWTGQSNHGIGYESDNHTAPILCGAQYRIHTHGVFRGIQDV
.....AAAAAAA.....AAA.AAA.....AAAAAAA.....
.....RRRRRRRRR.....RRRR.....
.....DDDDDDDDDD.....ddddd.....

305 310 315 320 325 330 335 340 345 350 355 360 365 370 375
RRVSGVAPTLVRSASETSEKRPFMCAYPGCNKRYFKLSHLQMHSRKHTGEKPYQCDFKDCERRFSRSRSDLQLKRHQR
.....AAAAAAA.....AAAAA.....AAAAA.....AAAAA.....
.....RRRR.....
.....DDDDDDDDDD.....

380 385 390 395 400 405 410 415 420 425 430 435 440 445 450
RHTGVKPFQCKTCORKFSRSDLHKTHTRHTGKTSEKPFSCRWHSCQKKFARSDELVRHHNMQRNMTKLHVAL
.....AAAA.AAA..AA.....AAAA.....AA.....AAAAAAA..AAA.....
.....RRRR.....RRRR.....
.....ddddd.....

FIG. 8B